

Supplemental Data

On the Use of General Control Samples

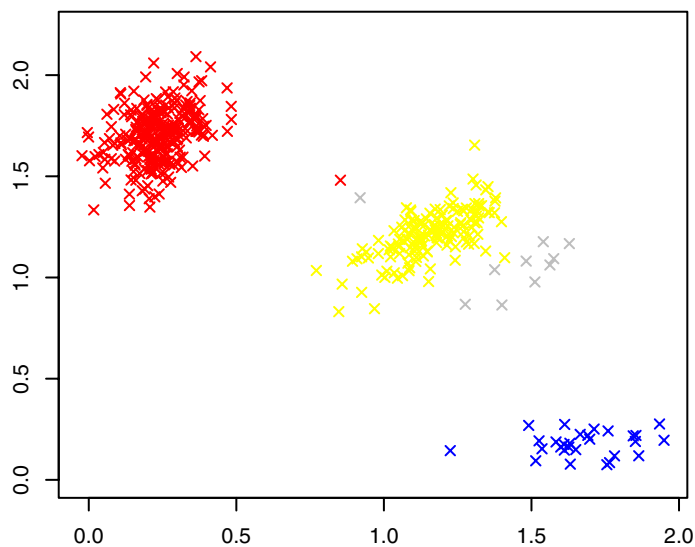
for Genome-wide Association Studies:

Genetic Matching Highlights Causal Variants

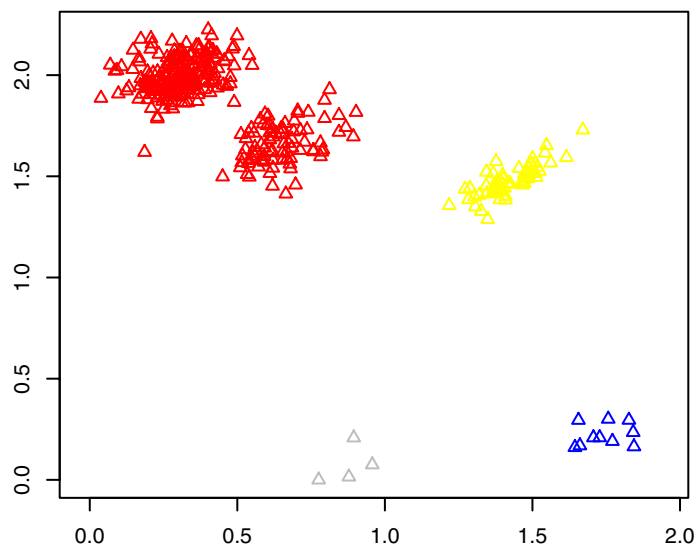
Diana Luca, Steven Ringquist, Lambertus Klei, Ann B. Lee, Christian Gieger, H.-Erich Wichmann, Stefan Schreiber, Michael Krawczak, Ying Lu, Alexis Styche, Bernie Devlin, Kathryn Roeder, and Massimo Trucco

Figure S1.Genotype Cluster Plots for All SNPs Producing a p Value < 0.00001 for the fMatch Analysis

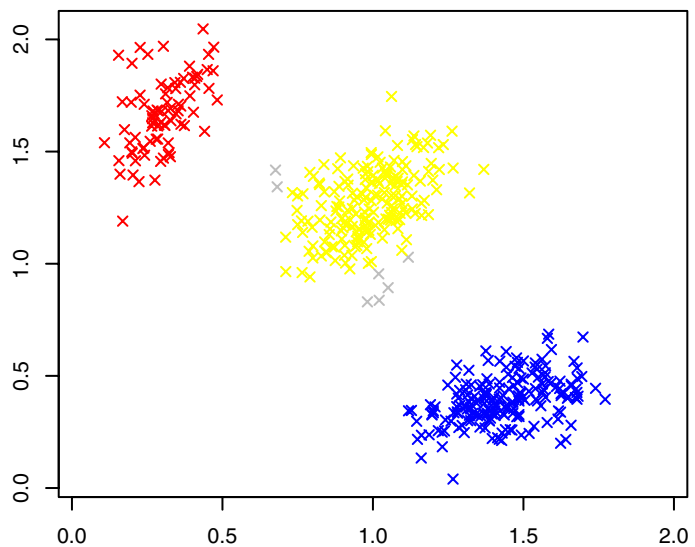
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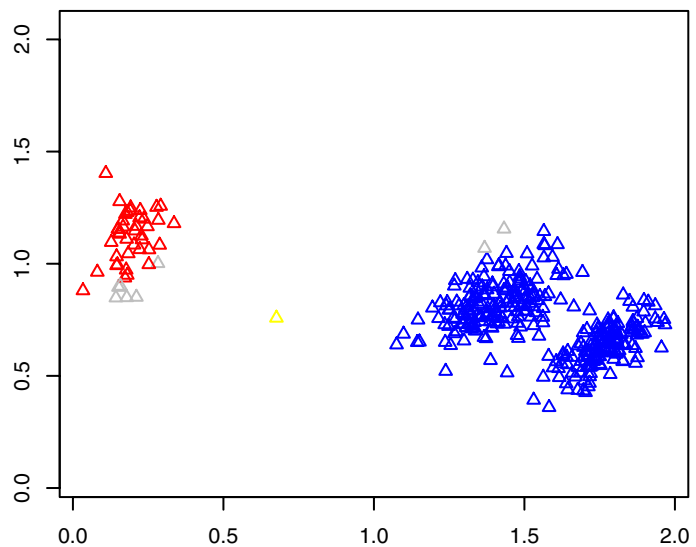
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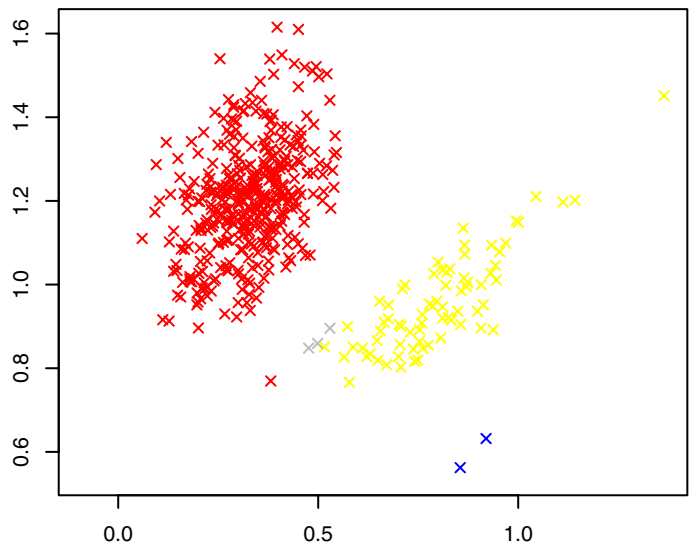
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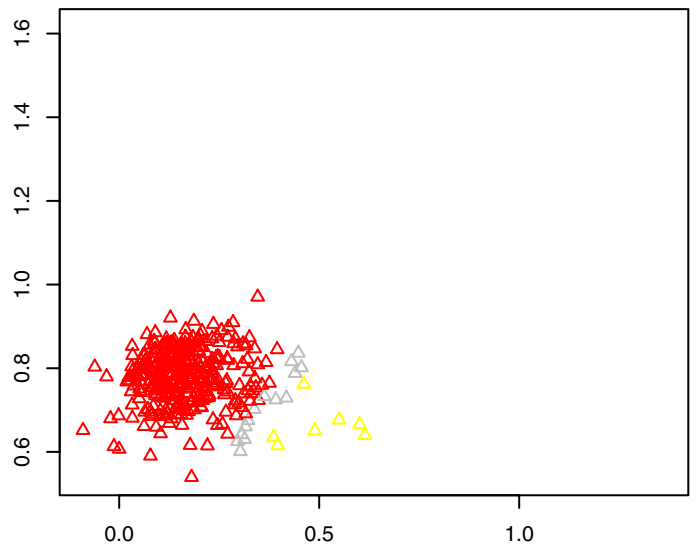
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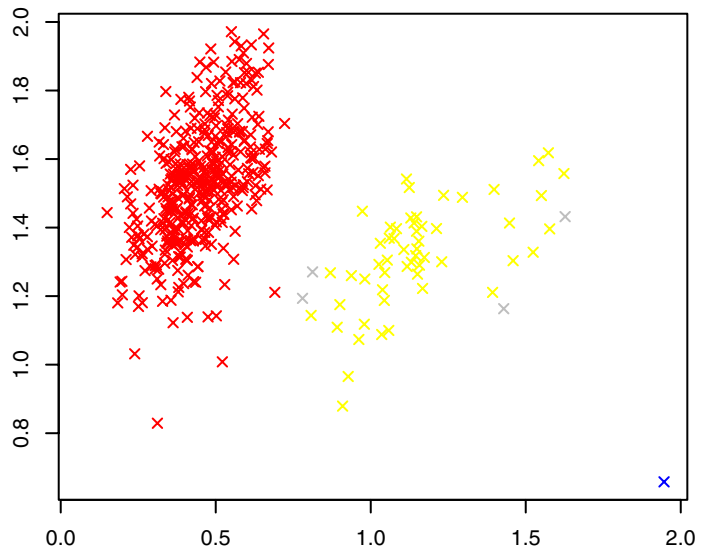
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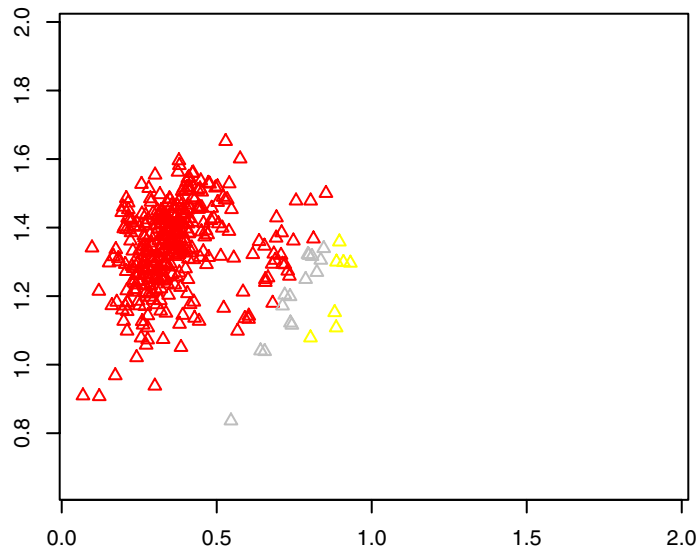
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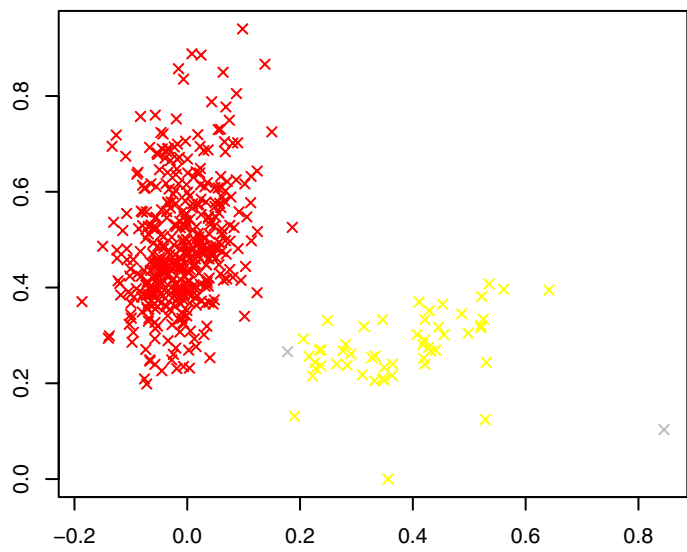
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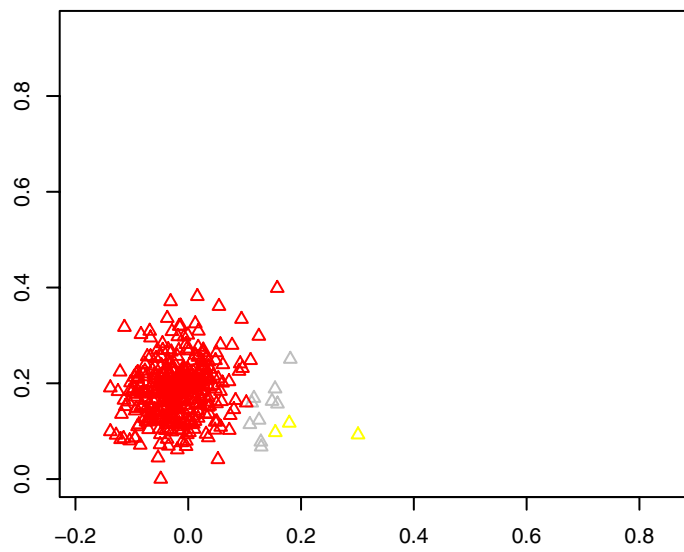
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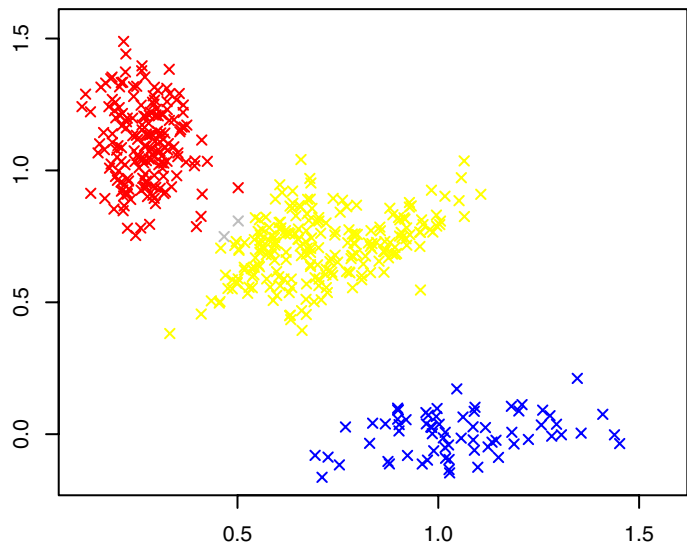
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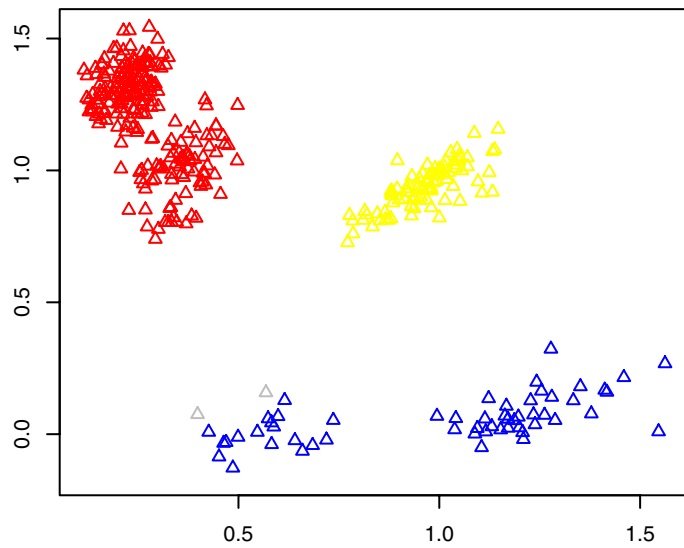
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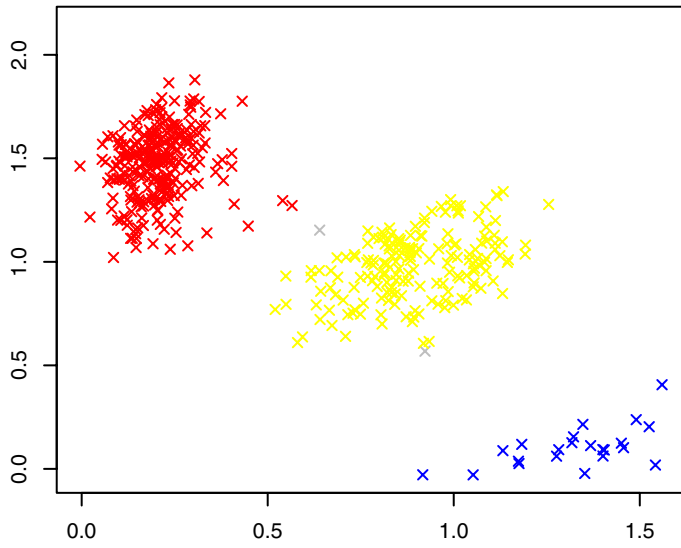
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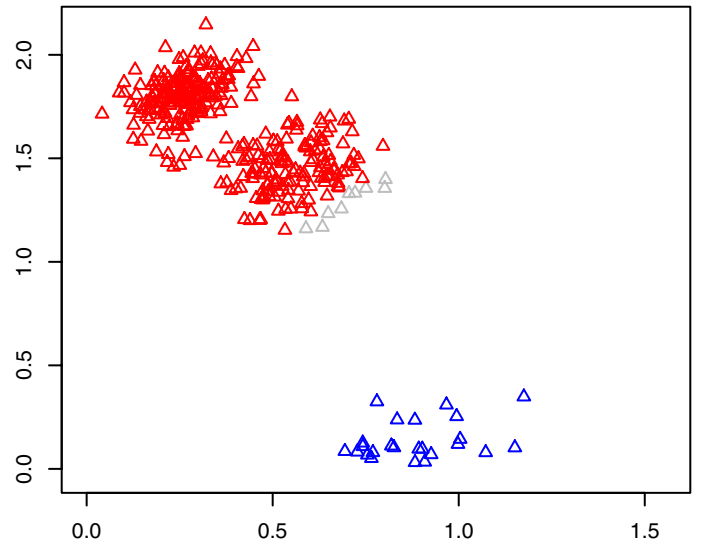
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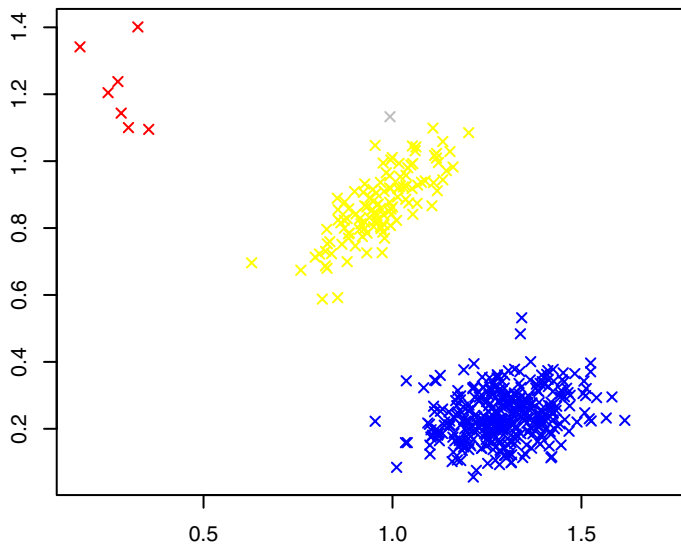
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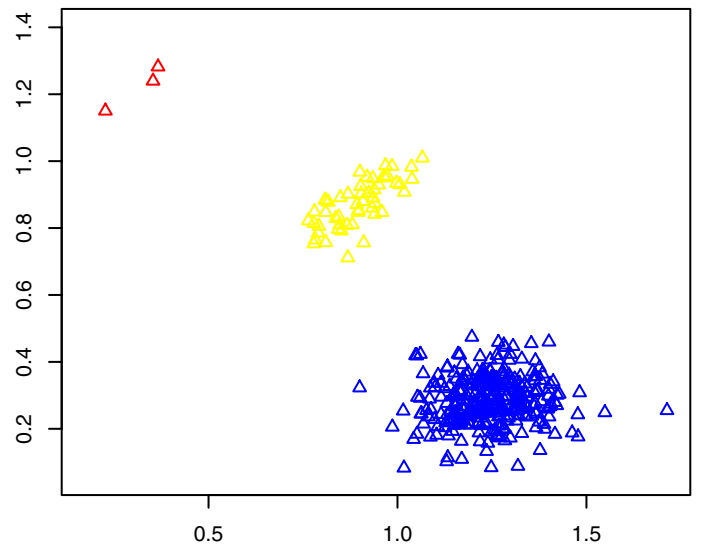
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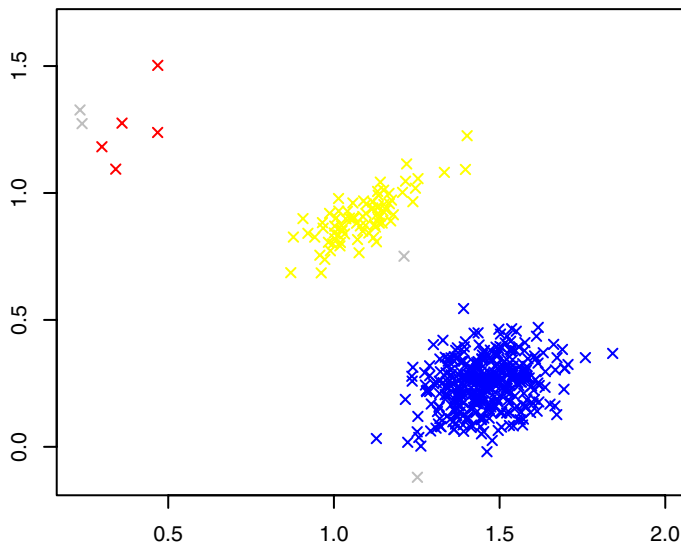
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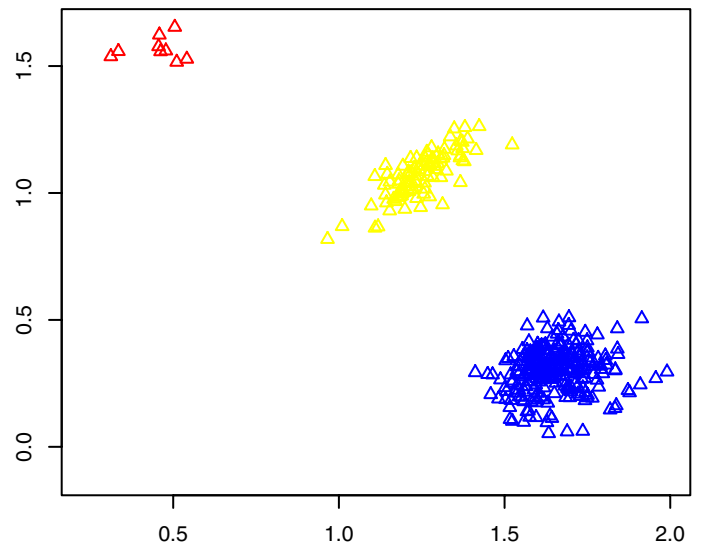
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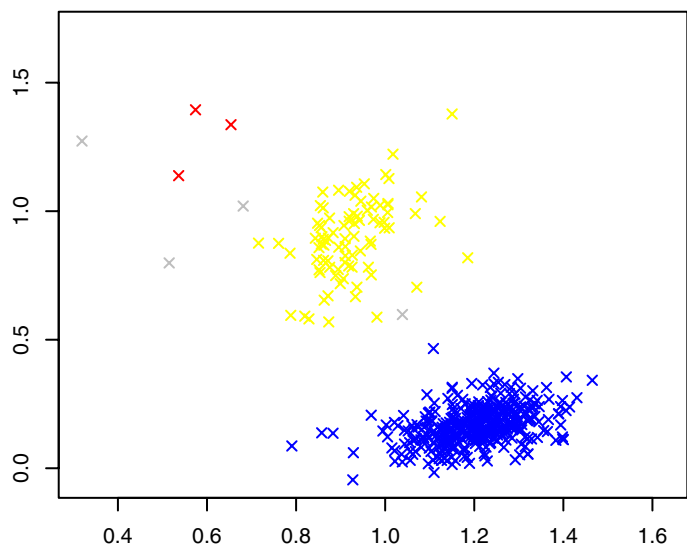
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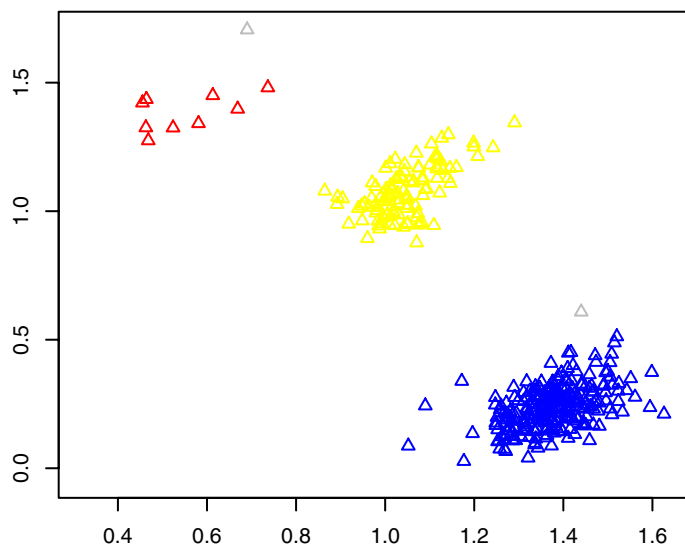
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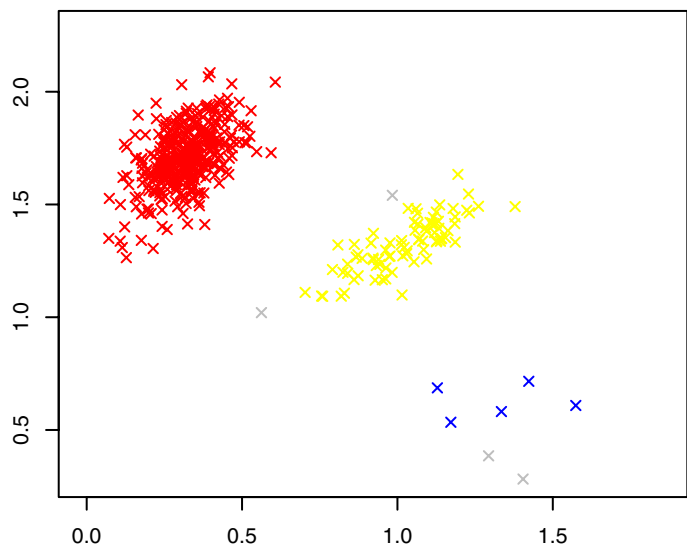
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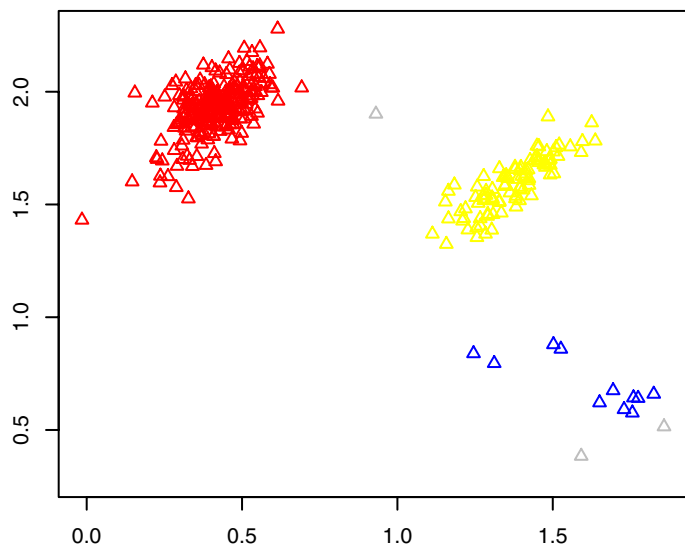
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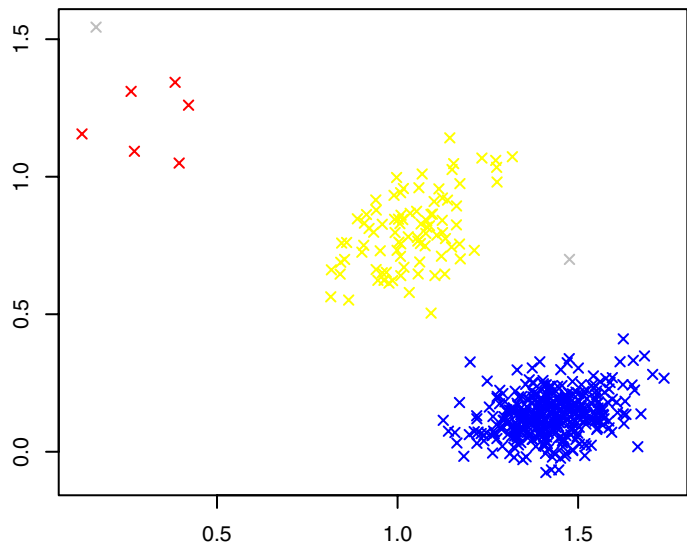
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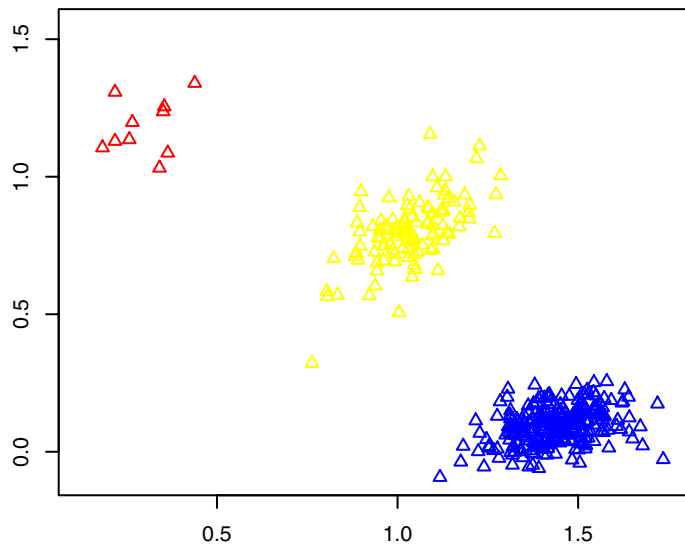
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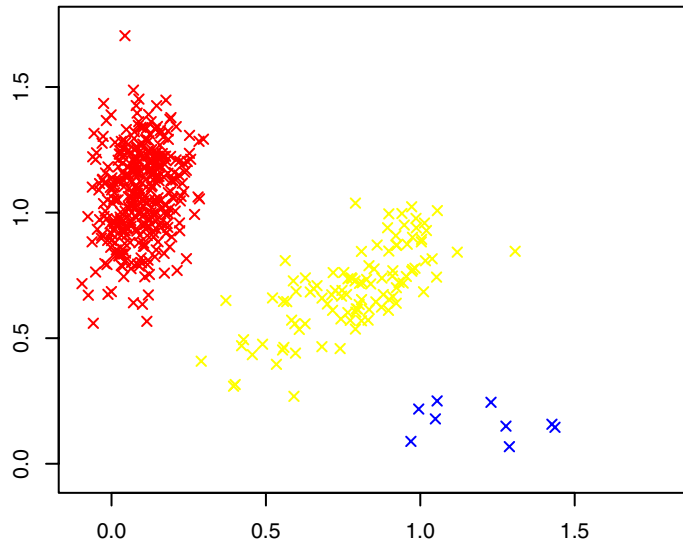
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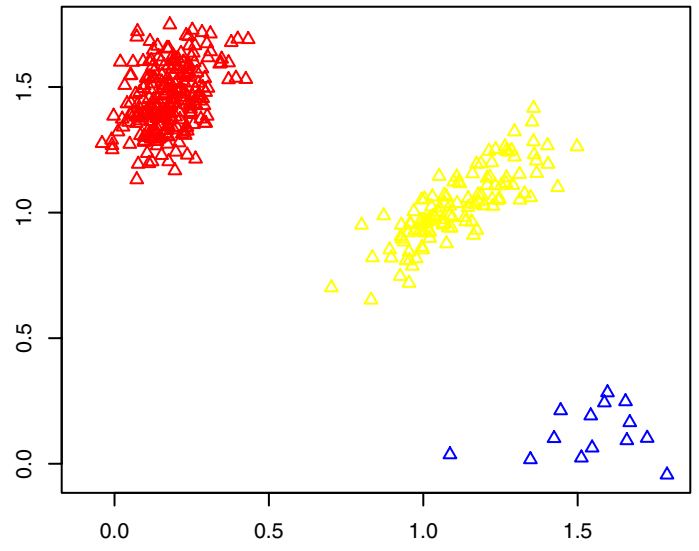
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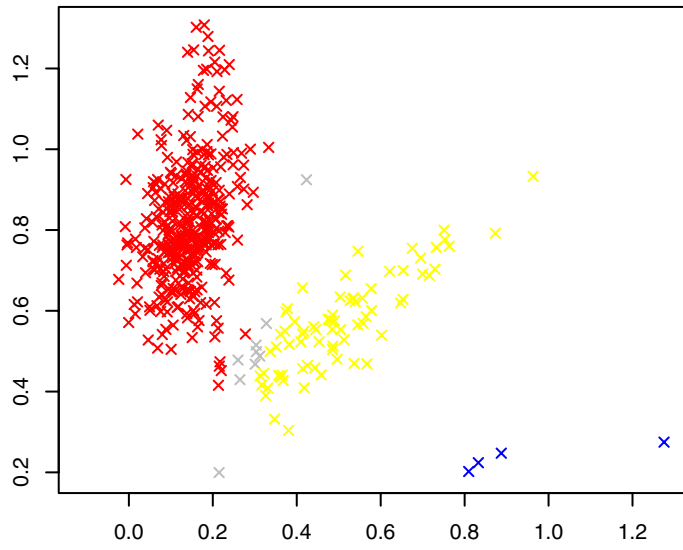
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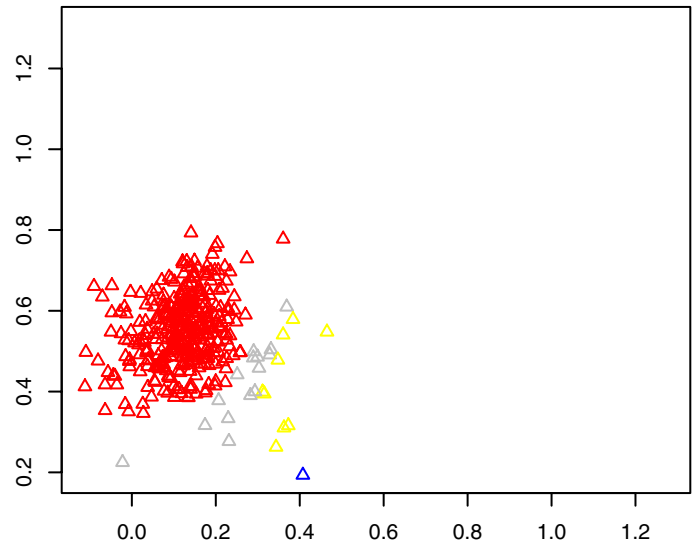
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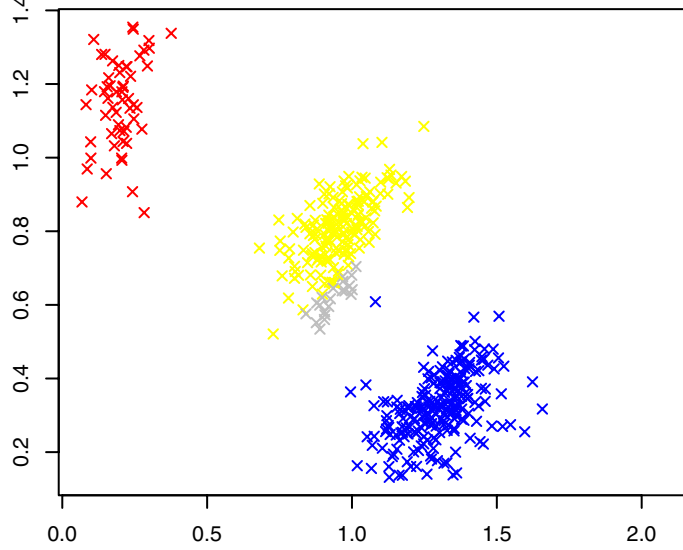
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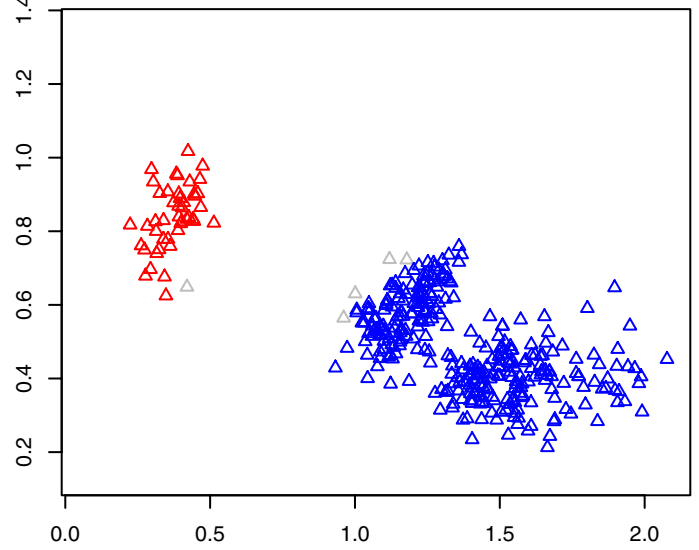
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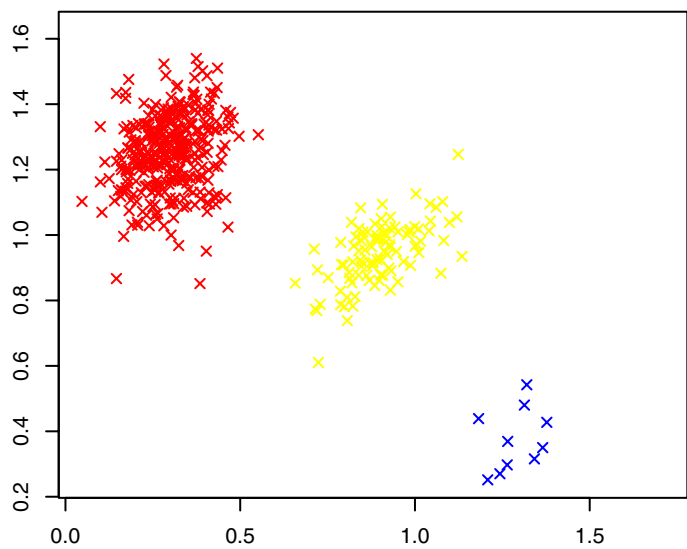
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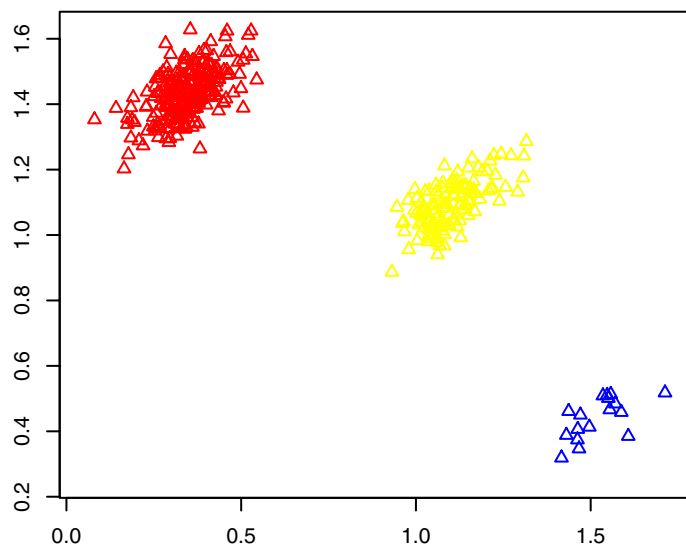
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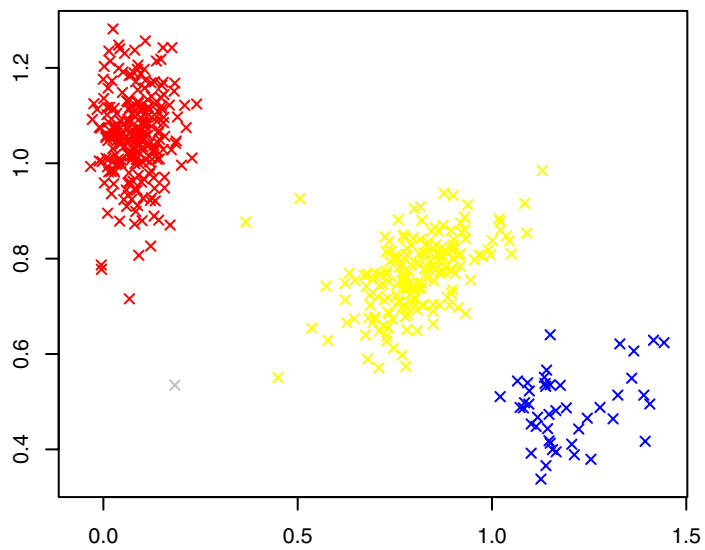
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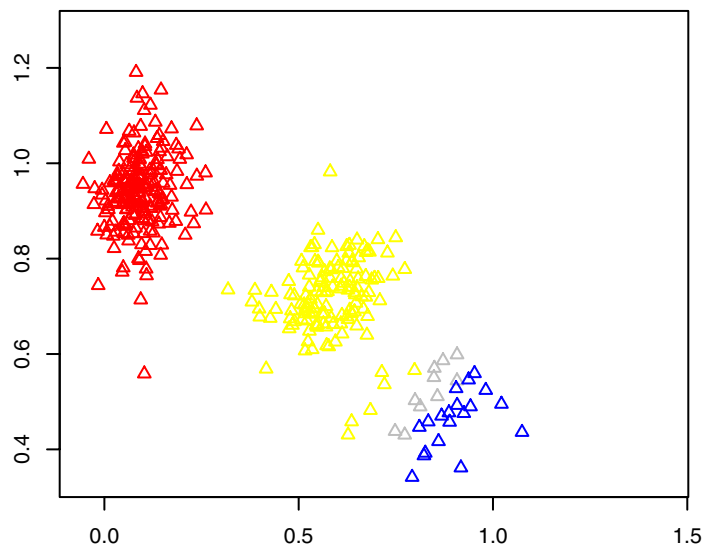
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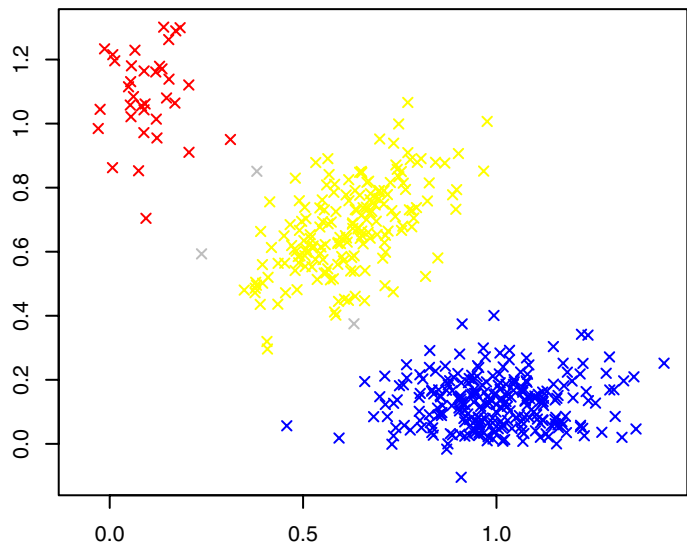
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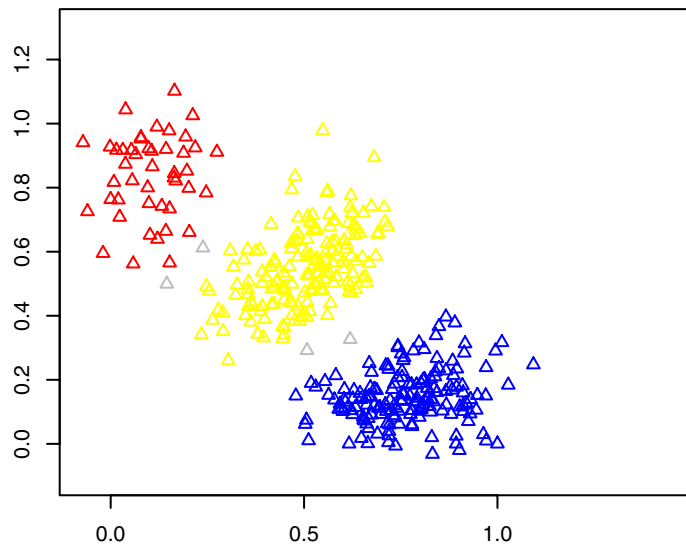
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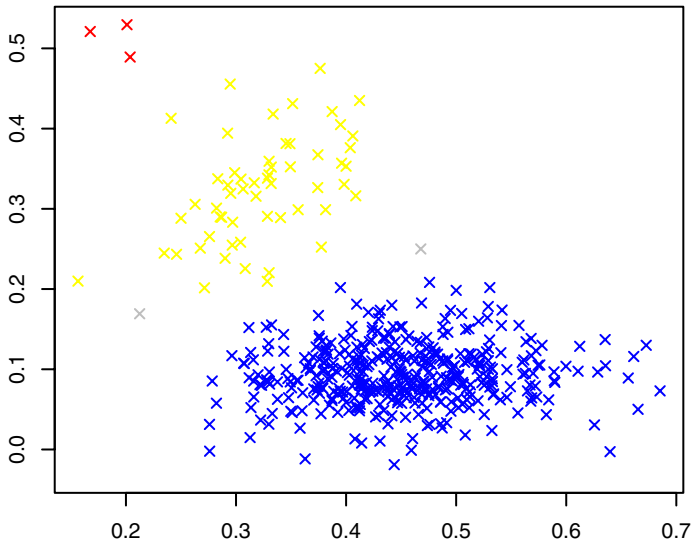
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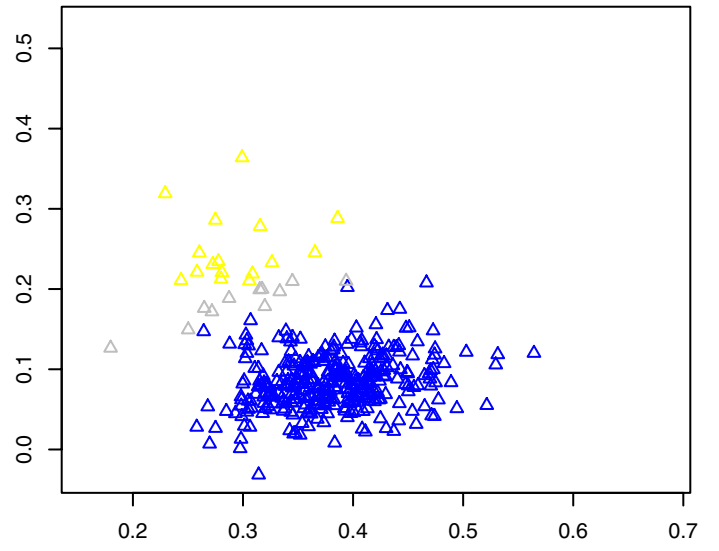
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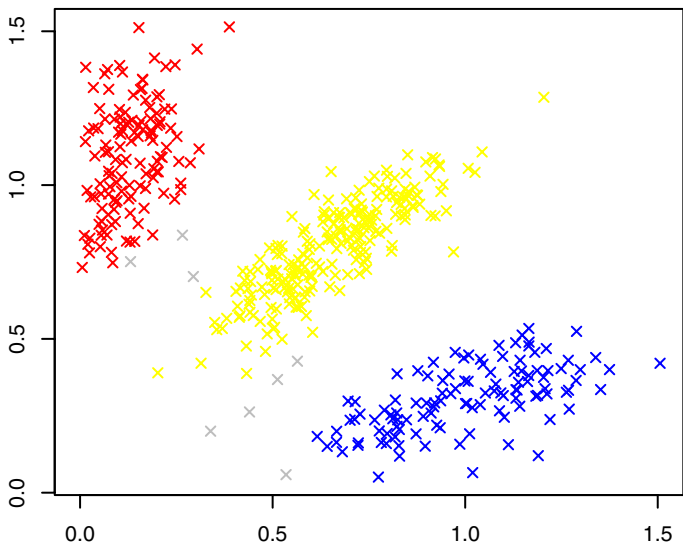
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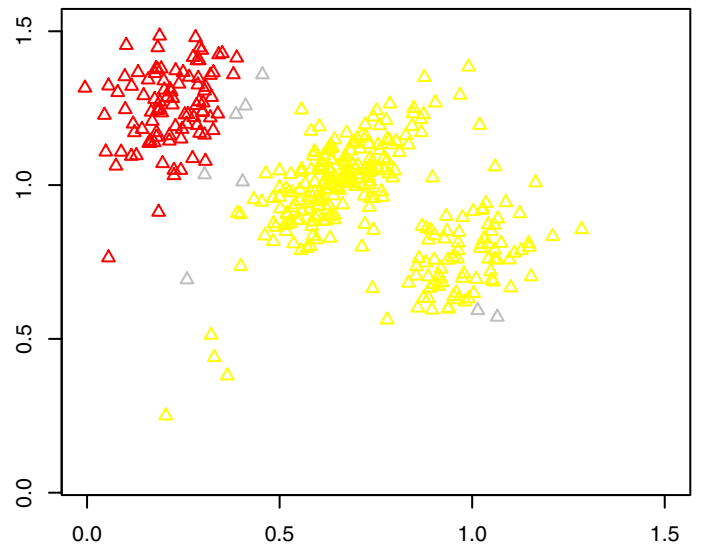
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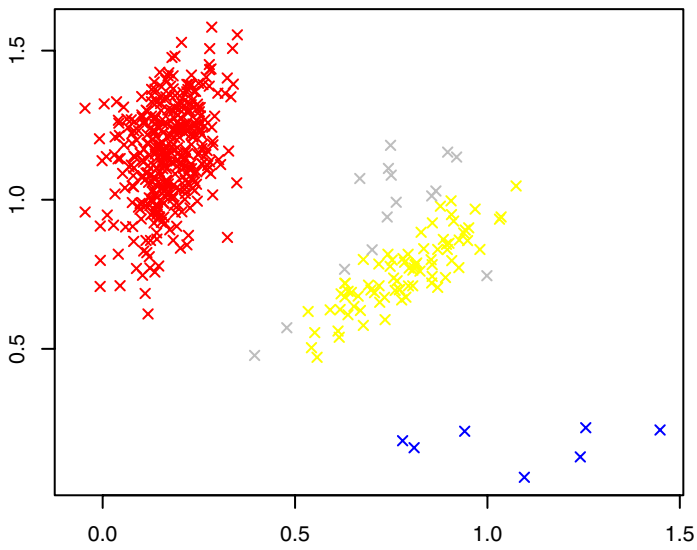
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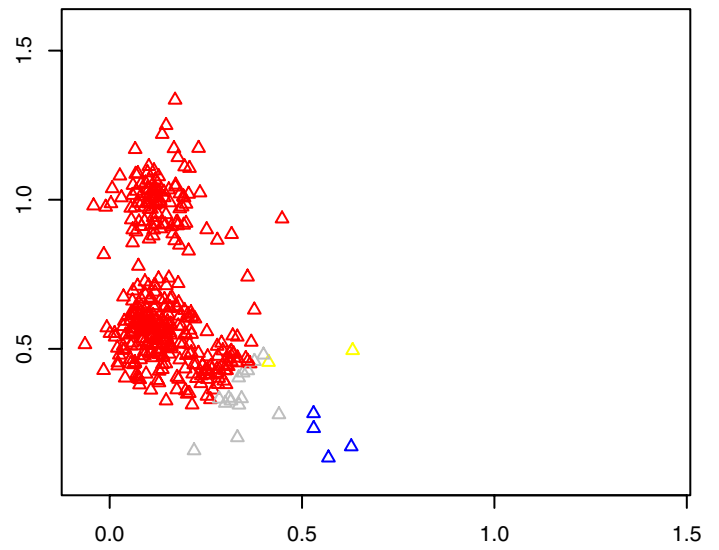
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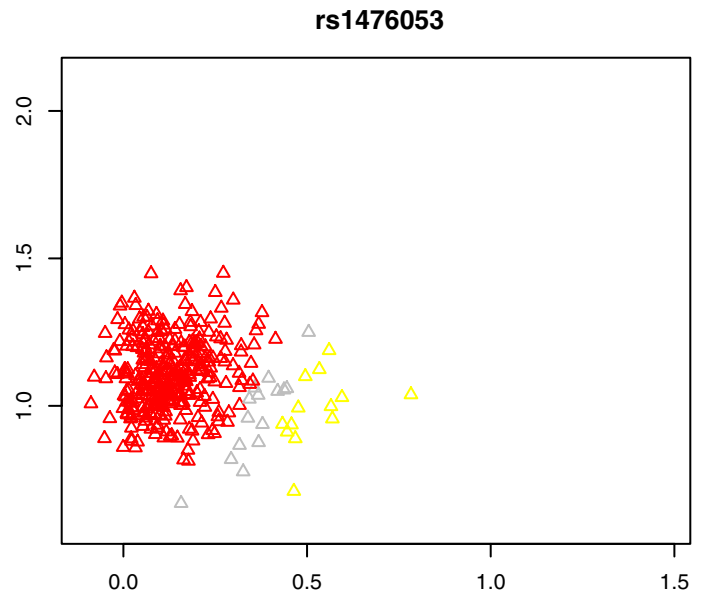
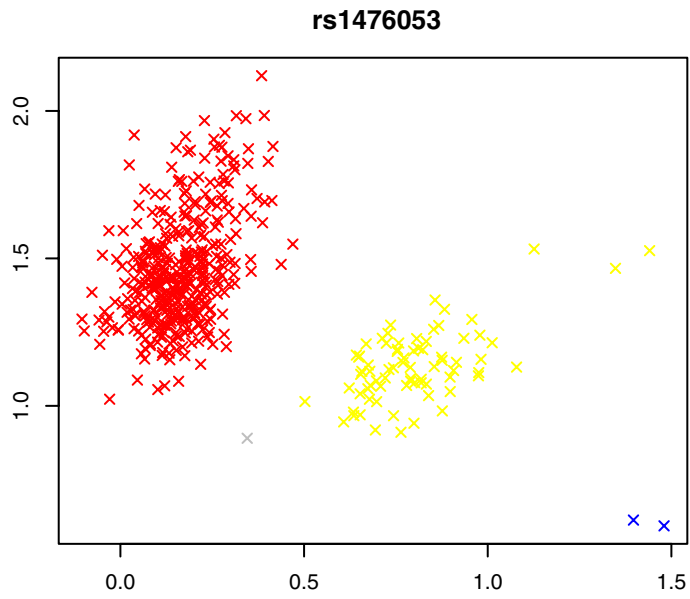


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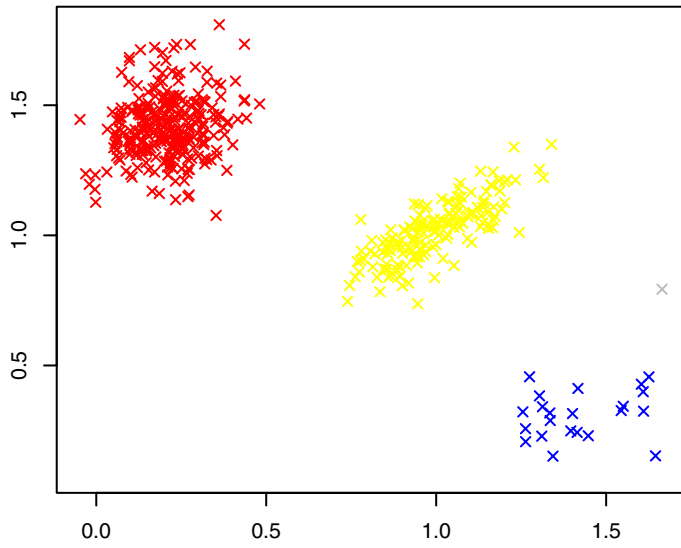




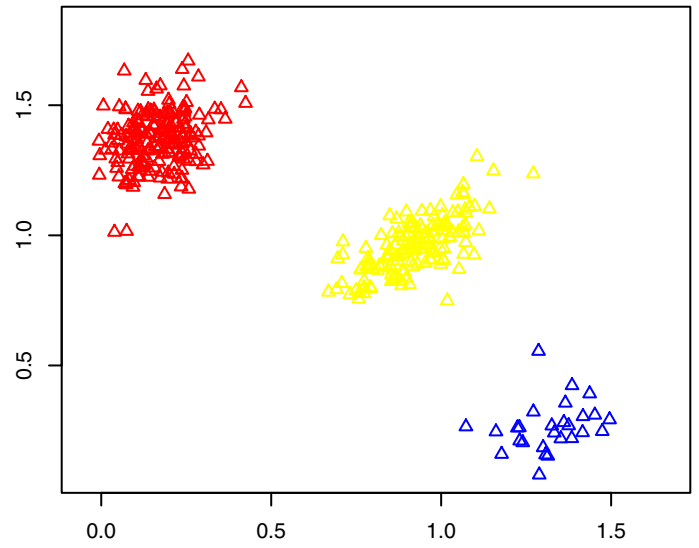
Previous findings for T1D risk loci are from ²⁸ and ²⁹. Left column shows results from controls; right shows results from cases. Colors highlight the three genotype clusters, with gray being uncalled genotypes.

Figure S2. Genotype Cluster Plots for All SNPs Cited in the Manuscript Supporting Previous Findings for T1D Risk Loci.²⁸⁻²⁹

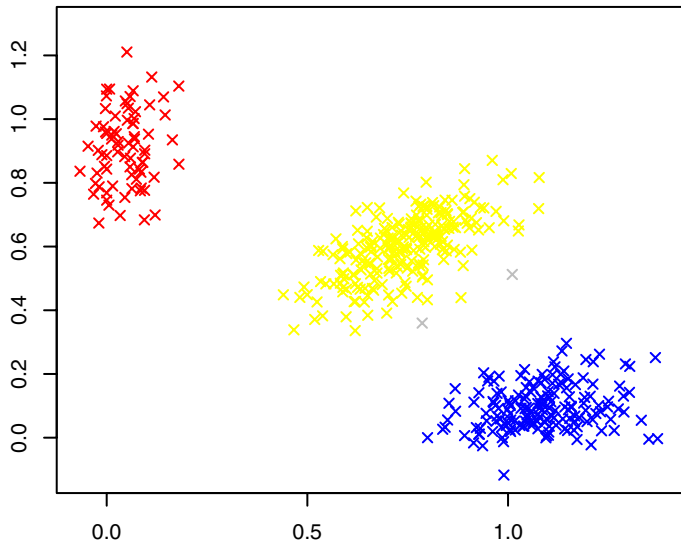
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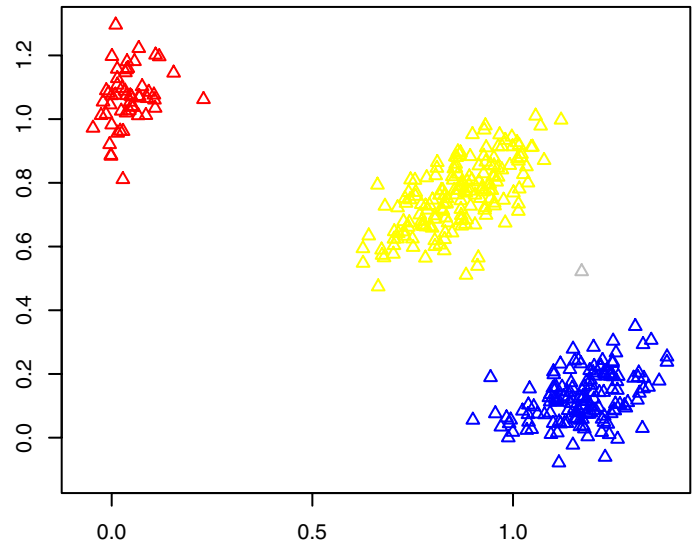
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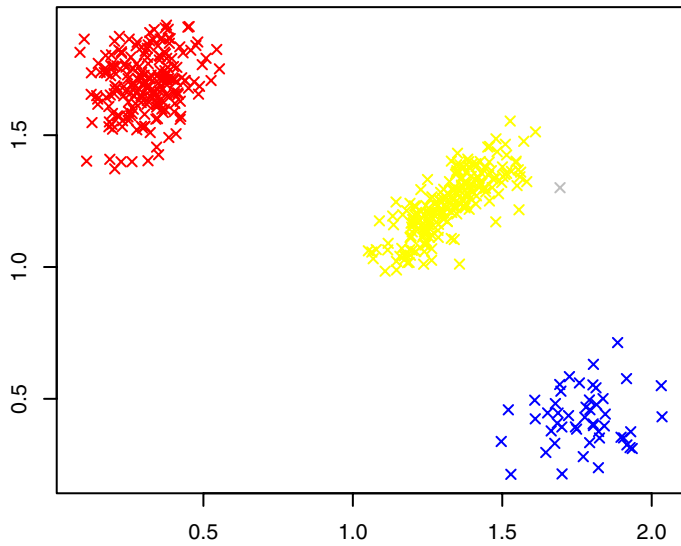
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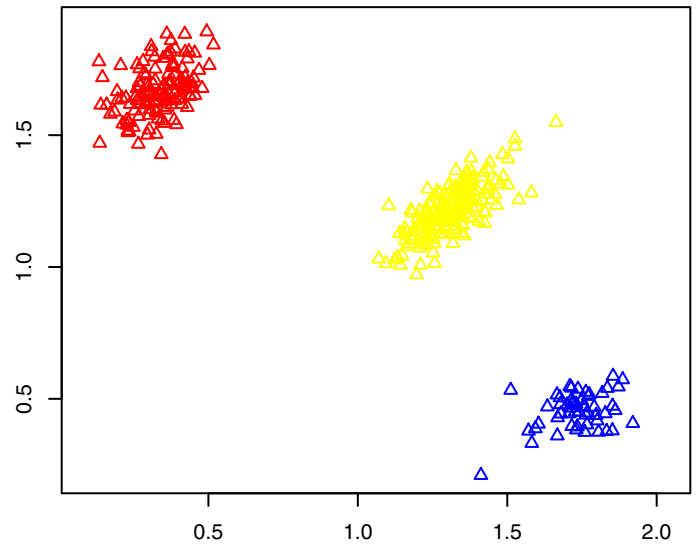
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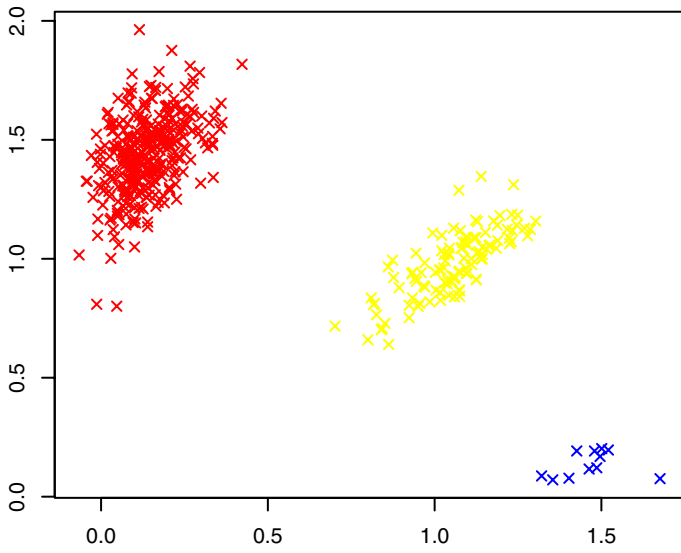
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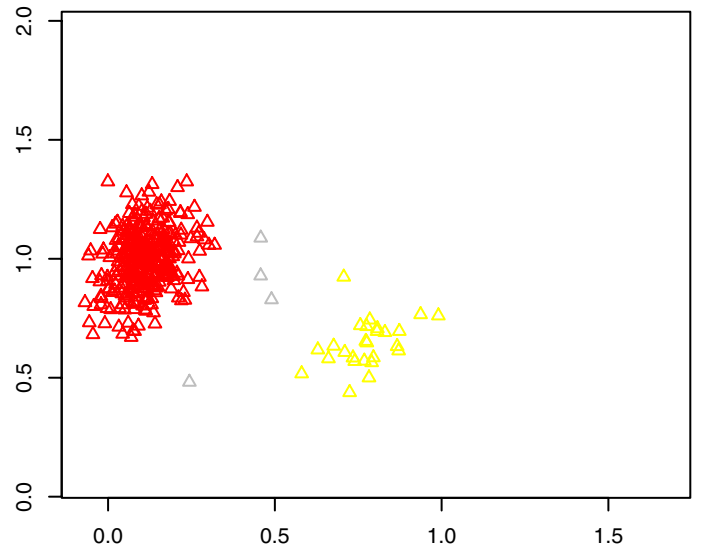
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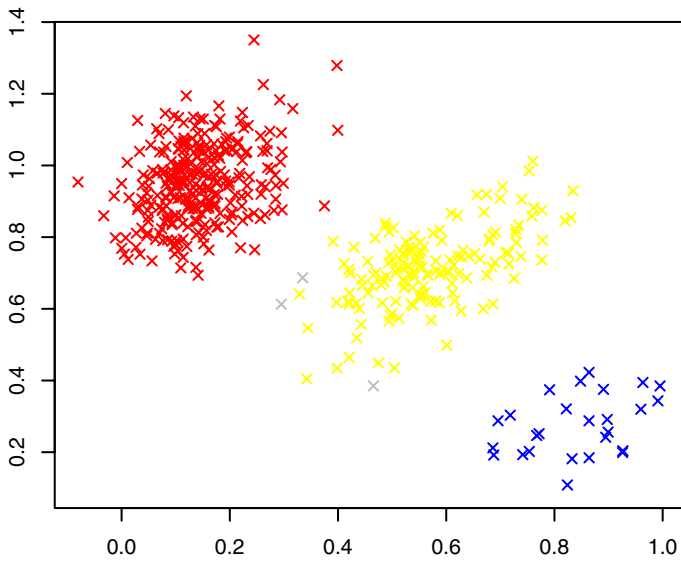
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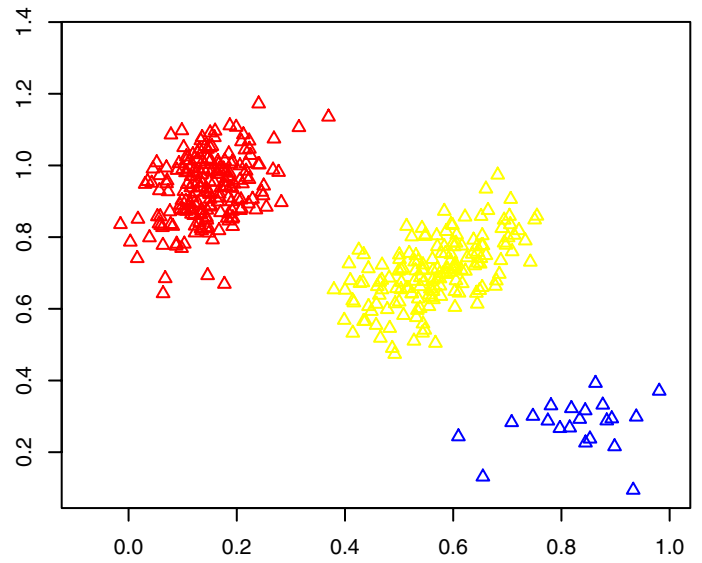
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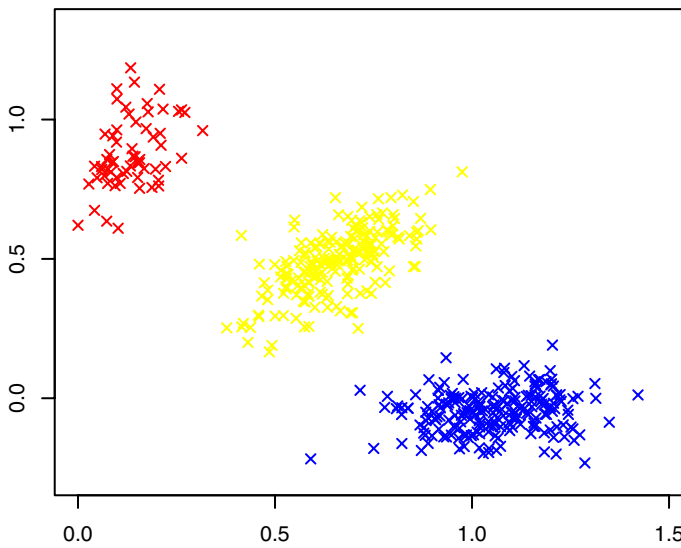
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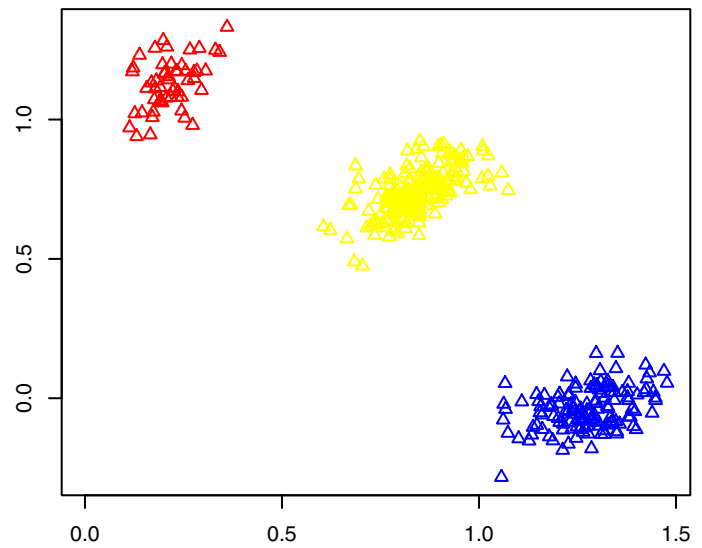
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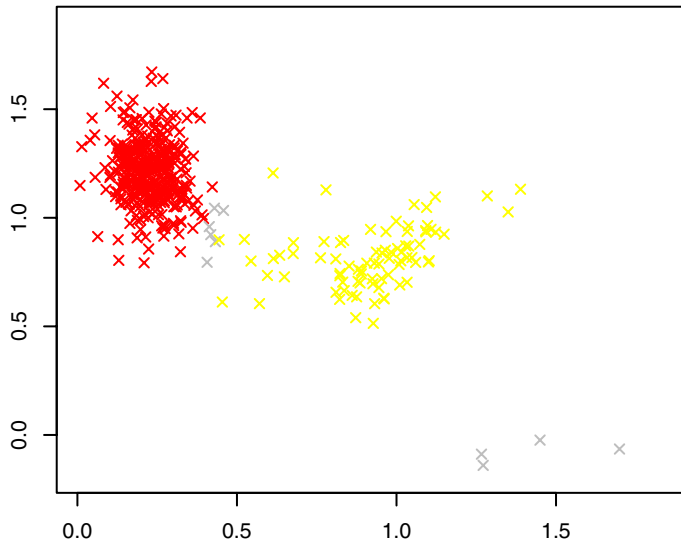
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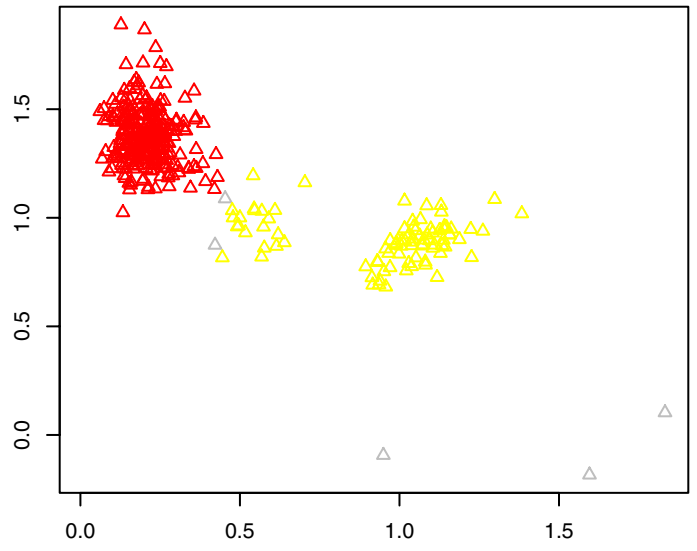
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rs11647011



The left column shows results from controls; the right shows results from cases. Colors highlight the three genotype clusters, with gray being uncalled genotypes.